



H16

1

SEQUENCE LISTING

<110> DENNIS, EDWARD  
WANG, AIJUN

<120> CLONED HUMAN LYSOPHOSPHOLIPASE

<130> 041673-0301

<140> 09/493,601  
<141> 2000-01-28

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<170> PatentIn Ver. 3.2

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<212> DNA

<213> Homo sapiens

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aaaacctatg aaggtatgtat gcacagttcg tgtcaacagg aaatgatgga tgtcaagccaa 660  
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<212> PRT

<213> Homo sapiens

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Thr Gly His Gly Trp Ala Glu Ala Phe Ala Gly Ile Arg Ser Ser His  
35 40 45

Ile Lys Tyr Ile Cys Pro His Ala Pro Val Arg Pro Val Thr Leu Asn  
50 55 60

Met Asn Val Ala Met Pro Ser Trp Phe Asp Ile Ile Gly Leu Ser Pro  
65 70 75 80

Asp	Ser	Gln	Glu	Asp	Glu	Ser	Gly	Ile	Lys	Gln	Ala	Ala	Glu	Asn	Ile
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Lys	Ala	Leu	Ile	Asp	Gln	Glu	Val	Lys	Asn	Gly	Ile	Pro	Ser	Asn	Arg
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Ile	Ile	Leu	Gly	Gly	Phe	Ser	Gln	Gly	Gly	Ala	Leu	Ser	Leu	Tyr	Thr
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Ala	Leu	Thr	Thr	Gln	Gln	Lys	Leu	Ala	Gly	Val	Thr	Ala	Leu	Ser	Cys
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Trp	Leu	Pro	Leu	Arg	Ala	Ser	Leu	Pro	Gln	Gly	Pro	Ile	Gly	Gly	Ala
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Asn	Arg	Asp	Ile	Ser	Ile	Leu	Gln	Cys	His	Gly	Asp	Cys	Asp	Pro	Leu
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Val	Pro	Leu	Met	Phe	Gly	Ser	Leu	Thr	Val	Glu	Lys	Leu	Lys	Thr	Leu
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Val	Asn	Pro	Ala	Asn	Val	Thr	Phe	Lys	Thr	Tyr	Glu	Gly	Met	Met	His
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Ser	Ser	Cys	Gln	Gln	Glu	Met	Met	Asp	Val	Lys	Gln	Phe	Ile	Asp	Lys
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Asn Asn Met Ser Thr Pro Leu Pro Ala Ile Val Pro Ala Ala Arg Lys
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gcc acc gct gcg gtg att ttc ctg cat gga ttg gga gat act ggg cac      153
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gag gat gaa tct ggg att aaa cag gca gca gaa aat ata aaa gct ttg Glu Asp Glu Ser Gly Ile Lys Gln Ala Ala Glu Asn Ile Lys Ala Leu 85 90 95	345
att gat caa gaa gtg aag aat ggc att cct tct aac aga att att ttg Ile Asp Gln Glu Val Lys Asn Gly Ile Pro Ser Asn Arg Ile Ile Leu 100 105 110 115	393
gga ggg ttt tct cag gga gga gct tta tct tta tat act gcc ctt acc Gly Gly Phe Ser Gln Gly Gly Ala Leu Ser Leu Tyr Thr Ala Leu Thr 120 125 130	441
aca cag cag aaa ctg gca ggt gtc act gca ctc agt tgc tgg ctt cca Thr Gln Gln Lys Leu Ala Gly Val Thr Ala Leu Ser Cys Trp Leu Pro 135 140 145	489
ctt cgg gct tcc ctt cca cag ggt cct atc ggt ggt gct aat aga gat Leu Arg Ala Ser Leu Pro Gln Gly Pro Ile Gly Gly Ala Asn Arg Asp 150 155 160	537
att tct att ctc cag tgc cac ggg gat tgt gac cct ttg gtt ccc ctg Ile Ser Ile Leu Gln Cys His Gly Asp Cys Asp Pro Leu Val Pro Leu 165 170 175	585
atg ttt ggt tct ctt acg gtg gaa aaa cta aaa aca ttg gtg aat cca Met Phe Gly Ser Leu Thr Val Glu Lys Leu Lys Thr Leu Val Asn Pro 180 185 190 195	633
gcc aat gtg acc ttt aaa acc tat gaa ggt atg atg cac agt tcg tgt Ala Asn Val Thr Phe Lys Thr Tyr Glu Gly Met Met His Ser Ser Cys 200 205 210	681
caa cag gaa atg atg gat gtc aag caa ttc att gat aaa ctc cta cct Gln Gln Glu Met Met Asp Val Lys Gln Phe Ile Asp Lys Leu Leu Pro 215 220 225	729
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